PC-0039 US

<110> Chen, Huei-Mei
      Honchell, Cynthia D.
      Tang, Y. Tom

<120> Mucin-Related Tumor Marker

<130> PC-0039 US

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Ser Tyr Ile Lys Ile Ser Asn Ser Ser His Ser Glu Tyr Ser Ser
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Phe Ser His Ala Gln Thr Glu Arg Ser Asn Ile Ser Ser Tyr Asp
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Gly Glu Tyr Ala Gln Pro Ser Thr Glu Ser Pro Val Leu His Thr
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Ser Val Val Leu Asp Thr Asp Ala Glu Phe Val Ser Asp Ser Ser
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PC-0039 US

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agacttggct cccaaatctg ccacctttgc tgttcagagc agcacacagt caccaacaac 180
actgtcctct tcagcctcag tcaacagctg tgctgtgaac ccttgtcttc acaatggcga 240
atgcgtcgca gacaacacca gccgtggcta ccactgcagg tgcccgcctt cctggcaagg 300
ggatgattgc agtgtggatg tgaatgagtg cctgtcgaac ccctgcccat ccacagccac 360
gtgcaacaat actcagggat cctttatctg caaatgcccg gttgggtacc agttggaaaa 420
agggatatgc aatttggtta gaaccttcgt gacagagttt aaattaaaga gaactttct 480
taatacaact gtggaaaaac attcagacct acaagaagtt gaaaatgaga tcaccaaaac 540
gttaaatatg tgtt                                             554


<210> 7
<211> 571
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 71247228V1

<400> 7
gatcaccaaa acgttaaata tgtgttttttc agcgttacct agttacatcc gatctacagt 60
tcacgcctct agggagtcca acgcggtggt gatctcactg caaacaacct tttccctggc 120
```

8

```
ctccaatgtg acgctatttg acctggctga taggatgcag aaatgtgtca actcctgcaa 180
ggtcctctgc tgaggtctgc cagctcttgg gatctcagag gcggatcttt agagcgggca 240
gcttgtgcaa gcggaagagt cccgaatgtg acaaagacac ctccatctgc actgacctgg 300
acggcgttgc cctgtgccag tgcaagtcgg gatactttca gttcaacaag atggaccact 360
cctgccgagc atgtgaagat ggatataggc ttgaaaatga aacctgcatg agttgcccat 420
ttggccttgg tggtctcaac tgtggaaacc cctatcagct tatcactgtg gtgatcgcag 480
ccgcgggagg tgggctcctg ctcatcctag gcatcgcact gattgttacc tgttgcagaa 540
agaataaaaa tgacataagc aaactcatct t                               571
```

```
<210> 8
<211> 433
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 6475676H1

<400> 8
tgaaacttgc atgagttgtc cattcagcct tggtggtctc aactgtggaa acccctatca 60
gcttatcact gtggtgatcg cagccgcggg aggtgggctc ctgctcatcc taggcatcgc 120
actgattgtt acctgttgca gaaagaataa aaatgacata agcaaactca tcttcaaaag 180
tggagatttc caaatgtccc cgtatgctga ataccccaaa aatcctcgct cacaagaatg 240
gggccgagaa gctattgaaa tgcatgagaa tggaagtacc aaaaacctcc tccagatgac 300
ggatgtgtac tactcgccta caagtgtaag gaatccagaa cttgaacgaa acggactcta 360
cccgggctac actggactgc caggatcacg ggattcttgc attttccccg gacagtataa 420
accgtctttc atc                                                   433
```

```
<210> 9
<211> 538
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 7735769H1

<400> 9
ggggccgaga agctattgaa atgcatgaga atggaagtac caaaaacctc ctccagatga 60
cggatgtgta ctactcgcct acaagtgtaa ggaatccaga acttgaacga acggactct 120
acccggccta cactggactg ccaggatcac ggcattcttg cattttcccc ggacagtata 180
acccgtcttt catcagtgat gaaagcagaa gaagagacta cttttaagtc caggagagag 240
agggactcat tgctctgagc cagtcacctg ggacctctgc tcagaggacc gcaccaggag 300
gctgcgccca ggatttgtcg ggagccacgc tgagtggcaa gcaggaacga gggacaggca 360
tgcggggcgt gaccacagtg gaggagacag gtggatgtgg aaccacaggc tgctcattca 420
gcacctttgt tgttactgtg aacgtgaatg tgggccagta tcaagagagt ctctctgagt 480
gactgcacca tggcactggc accagggcga ctattagcca gggcagacca ctagactt   538
```

```
<210> 10
<211> 567
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 7180688H1

<400> 10
ctagacttca gtgcaggacc tggttttccc ttcgtttgca ctttagtaaa ttgggtggga 60
ggtttccttt tggatctgtt ttgagactgt tccagaaaga aggcttcctt tcccgagaca 120
cttccatagg cagcaatttg gtgattcatt tgcagcaaaa tactggcttg ttaattattt 180
```

9

```
tcctgcccag cgcctgcgtg ctaaacaaca gatgaggatg agcgtaccac tgaagtctga 240
agatgtcgcc attgaacgga cagtgttttc atatgtttct aggttgtctt atgctacagt 300
ttccaagcca gcccccacag tgaggaaatg tgtgaggcac cgcacacaac tgcaatgtgt 360
tttttaagtc aaggtgacac atgtatttaa gatttttttt taaaatctct ttgcagttaa 420
atctcacttt ttcaaacaag cctggatcag ggcaaaacaa cttatatttg gttttagctg 480
gaggctcagc aggcagattg caggcagggg ggcacttttc atccatgaga ggccagcctg 540
gggcctggga ctctgatcac cattgtg 567
```

<210> 11
<211> 600
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 70650868V1

<400> 11
```
ctcacttcat ccaaaaccag gatgccccaa agaaggccaa taaacacagt tccccaggtg 60
gcaattaaag acaccgagta ctggatgtct ccctggcagg acccacatca caggcataat 120
aaataagatg agtggaactt ccttccccgaa ggtcaaccct cagttcctcg accaaccgga 180
agtcttcagt tctcccacac tgactggaag tataaccacg tttctggagg gtgcgacaca 240
gccatatgaa gaggtacaaa tgactgggtg agaaaaaaaa gttatttctt cagccgaata 300
aacagctttg agtggttgaa agtttacatg gggtttgtgg acatgagatt ctgggtacaa 360
agtgctgcag tagccggtga gcaaactcat gtgtggctcc atctcggctc cctgttcttc 420
ctcaggaatc cacacagctt cccaaagcac tgttgatgca ggaaatctaa cctggctatt 480
cagcccatcc ctctaaccac atccagctgc aggggctcaa caagctgctt tcctagagtg 540
gtgaaacctg cgttcagttt gacattttct cctccataag caggttgctc tggcctccac 600
```

<210> 12
<211> 371
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 2359874T6

<400> 12
```
gaagaaacaa ccatgcccgg ggcagaactg tccccagtgg ctgcaccatt tccagaaatg 60
tgaagcggga ctccccgctt cagggtggag acaattcttt tacctctgta ttcccctcac 120
ttcatccaaa accaggatgc cccaaagaag gccaataaac acagttcccc aggtggcaat 180
taaagacacc gagtactgga tgtctccctg gcaggaccca catcacaggc ataataaata 240
agatgagtgg aacttccttc ccgaagtcaa ccctcagttc ctcgaccaac cggaagtctt 300
cagttctccc acactgactg gaagtataac cacgtttctg gagggtgcga cacagccata 360
tgaaggaatt c 371
```

<210> 13
<211> 399
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 2359874R6

<400> 13
```
cttcatatgg ctgtgtcgca ccctccagaa acgtggttat acttccagtc agtgtgggag 60
aactgaagac ttccggttgg tcgaggaact gagggttgac cttcgggaag gaagttccac 120
tcatcttatt tattatgcct gtgatgtggg tcctgccagg gagacatcca gtactcggtg 180
tctttaattg ccacctgggg aactgtgttt attggccttc tttggggcat cctggttttg 240
```

```
gatgaagtga ggggaataca gaggtaaaag aattgtctcc accctgaagc ggggagtccc 300
gcttcacatt tctggaaatg gtgcagccac tggggacagt tctgccccgg gcatggttgt 360
ttcttcaagg tcctctaaat ataatcccta ttcttacat                        399
```

<210> 14
<211> 595
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 70650365V1

<400> 14
```
tttggggcat cctggttttg gatgaagtga ggggaataca gaggtaaaag aattgtctcc 60
accctgaagc ggggagtccc gcttcacatt tctggaaatg gtgcagccac tggggacagt 120
tctgccccgg gcatggttgt ttcttcaagg tcctctaaat ataatcccta ttcttacata 180
atcctgtggc ctgatggttt taagcaagaa ctcctgtgtc ccatggtctc caccactcac 240
catcaccctg ctgtagcaag agtcctagtc aggggaggtg cattttagta gttaaatggc 300
acttatccat gagataaata aaaggagaac tgtttttatc agtggaggct aacctaaaat 360
ttcaaagtgt cgccttttgg aaatctgggg cctctctctc tgtagaacca atggcccttg 420
gtggctcacg gcctcgcacc ctaactggag agttctgagc tcctgcagct cacctgagcc 480
cacagactag gcttcttggc tccttccgca gcaggctggt tcaccccaga acccgcagct 540
gtgggaagag ccatgtaggg aggctaatcc caggcataca cttccactgc cttca       595
```

<210> 15
<211> 549
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 1241344R6

<220>
<221> unsure
<222> 442, 460, 515
<223> a, t, c, g, or other

<400> 15
```
acctaactgg agagttctga gctcctgcag ctcacctgag cccacagact aggcttcttg 60
gctccttccg cagcatgcct gctcaccccc agaacccgca gctgtgggaa gagccatgta 120
gggaggctat tcccaggcat acacttccac tgccttcagc tgacgtcaca gctgacaaat 180
catctcctct atcggagcca gaagacttca gctccacaaa atgaagtgtt ctgtcctgaa 240
aacattcttg ggaagaatcc caacatcgag aaaacggtgt cctgtgagtt ccaacaatgc 300
ttcttgttca tgggtttctt ccgtatggag tggattaaga gtgttttatt ttgttgttct 360
aactgagaaa aaaaggaggc acccacaagg ttgaggtcac acagtctcca cagtttccag 420
gaggcgtttg ggggtgggga angcacctcc agagcatgan ggctctaagg ggacatgagt 480
aaagcatgtc tgtgacccag tgaggaaagg gagangccag ctgcactcct gcaacggggg 540
ttcctagct                                                        549
```

<210> 16
<211> 272
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 008938H1

<220>

11

<221> unsure
<222> 75, 106, 112, 163, 167, 192, 252
<223> a, t, c, g, or other

<400> 16
ggagaggcca gctgcactcc tgcacggggt tcctagctgc agaagggtcc cgcctaggcc 60
gaggggaaac acctnatagc agaagaggcc tggatgcaca cctggnacgc cnaggctctc 120
cgcccagaca cagtgctcca tgtcaacccc tgcacctggg gtntgtnatt cacgtgcaca 180
gatgccacaa tnctgcacca atatcccaca gatgggggaa ggtgagagga aggggcaagt 240
aatgtgtacc tnctcaagag atgcttaaac ct 272

<210> 17
<211> 424
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 2580841F6

<220>
<221> unsure
<222> 162, 251
<223> a, t, c, g, or other

<400> 17
ggtttaagct ccagagggtc ttattgccat tgtcttttcc tctgcccctt gagccagcct 60
aaggccctgg agtctgtttc tttaggcgga tgaactgaca tgctcctacc atgaccaggc 120
tctgggcaag gctcctcaca gtatccttga gaggtgggca tngaagtgcc catttctcag 180
gtacagaaac cttcagagag gataaatagc ttgccctgta gaagcaggac tgaaaccctt 240
gtccgcctga ntcccccagc tactctgccc actgtagccc cctgccttac tgtcctggca 300
cacccctcac catcctgtat accttaaata tcaaagaggg caagagagaa agggctttaa 360
agataagtta tttttttaag gaaccttaat attattttta agaagtaacc aaattagtga 420
cgtg 424

<210> 18
<211> 430
<212> DNA
<213> Homo sapiens

<220>
<221> misc_feature
<223> Incyte ID No: 70621193V1

<400> 18
cctggtacac ccctcaccat cctgtatacc ttaaatatca aagagggcaa gagagaaagg 60
gctttaaaga taagttattt ttttaaggaa ccttaatatt attttttaaga agtaaccaaa 120
ttagtgacgt gaaatgcaaa aaaaaaaaaa aaaaatgtct gactaccctt ttggaaaagt 180
gtgcttccag attggctttt ttatagtgta attctttaga cacttggtca ttaagaaaaa 240
tagtggcggg ctggtgcttc agcaagaagc acacggcac ggtggcttgg gatataggag 300
gtggaaggca aggaccgggt gtttctggac aggtggcggc cagacttaca cttccatctg 360
gagagctggt ggctttggtc ccctgggtag ggccatgggt tccccactat tactgggaag 420
ctatagggtg 430

<210> 19
<211> 957
<212> PRT
<213> Homo sapiens

<220>
<221> misc_feature

<223> Genbank ID No: g2853301

<400> 19

```
Ile Thr Ile Thr Glu Thr Thr Ser His Ser Thr Pro Ser Tyr Thr
1               5                   10                  15
Thr Ser Ile Thr Thr Thr Glu Thr Pro Ser His Ser Thr Pro Ser
                20                  25                  30
Tyr Thr Thr Ser Ile Thr Thr Thr Glu Thr Pro Ser His Ser Thr
                35                  40                  45
Pro Ser Phe Thr Ser Ser Ile Thr Thr Thr Glu Thr Thr Ser His
                50                  55                  60
Ser Thr Pro Ser Phe Thr Ser Ser Ile Arg Thr Thr Glu Thr Thr
                65                  70                  75
Ser Tyr Ser Thr Pro Ser Phe Thr Ser Ser Asn Thr Ile Thr Glu
                80                  85                  90
Thr Thr Ser His Ser Thr Pro Ser Tyr Ile Thr Ser Ile Thr Thr
                95                  100                 105
Thr Glu Thr Pro Ser Ser Ser Thr Pro Ser Phe Ser Ser Ser Ile
                110                 115                 120
Thr Thr Thr Glu Thr Thr Ser His Ser Thr Pro Gly Phe Thr Ser
                125                 130                 135
Ser Ile Thr Thr Thr Glu Thr Thr Ser His Ser Thr Pro Ser Phe
                140                 145                 150
Thr Ser Ser Ile Thr Thr Thr Glu Thr Thr Ser His Asp Thr Pro
                155                 160                 165
Ser Phe Thr Ser Ser Ile Thr Thr Ser Glu Thr Pro Ser His Ser
                170                 175                 180
Thr Pro Ser Ser Thr Ser Leu Ile Thr Thr Thr Lys Thr Thr Ser
                185                 190                 195
His Ser Thr Pro Ser Phe Thr Ser Ser Ile Thr Thr Thr Glu Thr
                200                 205                 210
Thr Ser His Ser Ala Arg Ser Phe Thr Ser Ser Ile Thr Thr Thr
                215                 220                 225
Glu Thr Thr Ser His Asn Thr Arg Ser Phe Thr Ser Ser Ile Thr
                230                 235                 240
Thr Thr Glu Thr Asn Ser His Ser Thr Thr Ser Phe Thr Ser Ser
                245                 250                 255
Ile Thr Thr Thr Glu Thr Thr Ser His Ser Thr Pro Ser Phe Ser
                260                 265                 270
Ser Ser Ile Thr Thr Thr Glu Thr Pro Leu His Ser Thr Pro Gly
                275                 280                 285
Leu Pro Ser Trp Val Thr Thr Thr Lys Thr Thr Ser His Ile Thr
                290                 295                 300
Pro Gly Leu Thr Ser Ser Ile Thr Thr Thr Glu Thr Thr Ser His
                305                 310                 315
Ser Thr Pro Gly Phe Thr Ser Ser Ile Thr Thr Thr Glu Thr Thr
                320                 325                 330
Ser Glu Ser Thr Pro Ser Leu Ser Ser Ser Thr Ile Tyr Ser Thr
                335                 340                 345
Val Ser Thr Ser Thr Thr Ala Ile Thr Ser His Phe Thr Thr Ser
                350                 355                 360
Glu Thr Ala Val Thr Pro Thr Pro Val Thr Pro Ser Ser Leu Ser
                365                 370                 375
Thr Asp Ile Pro Thr Thr Ser Leu Arg Thr Leu Thr Pro Ser Ser
                380                 385                 390
Val Gly Thr Ser Thr Ser Leu Thr Thr Thr Thr Asp Phe Pro Ser
                395                 400                 405
Ile Pro Thr Asp Ile Ser Thr Leu Pro Thr Arg Thr His Ile Ile
                410                 415                 420
Ser Ser Ser Pro Ser Ile Gln Ser Thr Glu Thr Ser Ser Leu Val
                425                 430                 435
```

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```
Gly Thr Thr Ser Pro Thr Met Ser Thr Val Arg Met Thr Leu Arg
            440             445             450
Ile Thr Glu Asn Thr Pro Ile Ser Ser Phe Ser Thr Ser Ile Val
            455             460             465
Val Ile Pro Glu Thr Pro Thr Gln Thr Pro Pro Val Leu Thr Ser
            470             475             480
Ala Thr Gly Thr Gln Thr Ser Pro Ala Pro Thr Thr Val Thr Phe
            485             490             495
Gly Ser Thr Asp Ser Ser Thr Ser Thr Leu His Thr Leu Thr Pro
            500             505             510
Ser Thr Ala Leu Ser Thr Ile Val Ser Thr Ser Gln Val Pro Ile
            515             520             525
Pro Ser Thr His Ser Ser Thr Leu Gln Thr Thr Pro Ser Thr Pro
            530             535             540
Ser Leu Gln Thr Ser Leu Thr Ser Thr Ser Glu Phe Thr Thr Glu
            545             550             555
Ser Phe Thr Arg Gly Ser Thr Ser Thr Asn Ala Ile Leu Thr Ser
            560             565             570
Phe Ser Thr Ile Ile Trp Ser Ser Thr Pro Thr Ile Ile Met Ser
            575             580             585
Ser Ser Pro Ser Ser Ala Ser Ile Thr Pro Val Phe Ser Thr Thr
            590             595             600
Ile His Ser Val Pro Ser Ser Pro Tyr Ile Phe Ser Thr Glu Asn
            605             610             615
Val Gly Ser Ala Ser Ile Thr Gly Phe Pro Ser Leu Ser Ser Ser
            620             625             630
Ala Thr Thr Ser Thr Ser Ser Thr Ser Ser Ser Leu Thr Thr Ala
            635             640             645
Leu Thr Glu Ile Thr Pro Phe Ser Tyr Ile Ser Leu Pro Ser Thr
            650             655             660
Thr Pro Cys Pro Gly Thr Ile Thr Ile Thr Ile Val Pro Ala Ser
            665             670             675
Pro Thr Asp Pro Cys Val Glu Met Asp Pro Ser Thr Glu Ala Thr
            680             685             690
Ser Pro Pro Thr Thr Pro Leu Thr Val Phe Pro Phe Thr Thr Glu
            695             700             705
Met Val Thr Cys Pro Thr Ser Ile Ser Ile Gln Thr Thr Leu Thr
            710             715             720
Thr Tyr Met Asp Thr Ser Ser Met Met Pro Glu Ser Glu Ser Ser
            725             730             735
Ile Ser Pro Asn Ala Ser Ser Ser Thr Gly Thr Gly Thr Val Pro
            740             745             750
Thr Asn Thr Val Phe Thr Ser Thr Arg Leu Pro Thr Ser Glu Thr
            755             760             765
Trp Leu Ser Asn Ser Ser Val Ile Pro Leu Pro Leu Pro Gly Val
            770             775             780
Ser Thr Ile Pro Leu Thr Met Lys Pro Ser Ser Ser Leu Pro Thr
            785             790             795
Ile Leu Arg Thr Ser Ser Lys Ser Thr His Pro Ser Pro Pro Thr
            800             805             810
Thr Arg Thr Ser Glu Thr Pro Val Ala Thr Thr Gln Thr Pro Thr
            815             820             825
Thr Leu Thr Ser Arg Arg Thr Thr Arg Ile Thr Ser Gln Met Thr
            830             835             840
Thr Gln Ser Thr Leu Thr Thr Thr Ala Gly Thr Cys Asp Asn Gly
            845             850             855
Gly Thr Trp Glu Gln Gly Gln Cys Ala Cys Leu Pro Gly Phe Ser
            860             865             870
Gly Asp Arg Cys Gln Leu Gln Thr Arg Cys Gln Asn Gly Gly Gln
            875             880             885
Trp Asp Gly Leu Lys Cys Gln Cys Pro Ser Thr Phe Tyr Gly Ser
```

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```
                          890                  895                   900
Ser Cys Glu Phe Ala Val Glu Gln Val Asp Leu Asp Ala Glu Asp
                          905                  910                   915
Phe Cys Arg His Ala Gly Leu His Leu Gln Gly Cys Gly Asp Pro
                          920                  925                   930
Val Pro Glu Glu Trp Gln His Arg Gly Gly Leu Pro Gly Pro Ala
                          935                  940                   945
Gly Asp Ala Leu Gln Pro Pro Ala Gly Glu Arg Val
                          950                  955
```

<210> 20
<211> 528
<212> PRT
<213> Sus scrofa

<220>
<221> misc_feature
<223> Genbank ID No: g915208

<400> 20
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Pro Ile Ser Val Gln Pro Ser Ser Ser Ser Ser Ser Pro Thr Thr
  1               5                  10                   15
Ser Thr Thr Ser Val Gln Ser Ser Ser Ser Ser Ser Val Pro Ile
                 20                  25                   30
Pro Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Gly Ser Ala Pro
                 35                  40                   45
Thr Thr Ser Ala Thr Ser Val Gln Thr Ser Ser Ser Ser Ser Pro
                 50                  55                   60
Pro Ile Ser Ser Thr Ile Ser Val Gln Thr Ser Ser Ser Ser Ser
                 65                  70                   75
Val Pro Thr Thr Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Ser
                 80                  85                   90
Ser Ala Pro Thr Thr Arg Ala Thr Ser Val Gln Ser Ser Ser Ser
                 95                  100                  105
Ser Ser Ala Pro Ile Ser Ser Thr Thr Ser Val Gln Pro Ser Ser
                 110                 115                  120
Ser Gly Ser Val Pro Thr Thr Ser Ala Thr Ser Val Gln Ser Ser
                 125                 130                  135
Ser Ser Ser Ser Ala Pro Thr Thr Ser Ala Thr Ser Val Gln Pro
                 140                 145                  150
Ser Ser Ser Ser Ser Pro Pro Ile Ser Ser Thr Val Ser Val Gln
                 155                 160                  165
Pro Ser Ser Ser Ser Ser Ala Pro Thr Thr Ser Ala Thr Ser Val
                 170                 175                  180
Gln Pro Ser Ser Ser Ser Ser Pro Pro Ile Ser Ser Thr Val Ser
                 185                 190                  195
Val Gln Thr Ser Ser Ser Ser Ser Val Pro Thr Thr Ser Thr Thr
                 200                 205                  210
Ser Val Gln Pro Ser Ser Ser Ser Ser Val Pro Thr Thr Ser Ala
                 215                 220                  225
Thr Ser Val Arg Ser Ser Ser Ser Ser Ser Thr Pro Ile Pro Ser
                 230                 235                  240
Thr Thr Ser Val Gln Pro Ser Ser Ser Ser Ala Pro Thr Thr
                 245                 250                  255
Ser Ala Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Thr Pro Ile
                 260                 265                  270
Pro Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Ser Ala Pro
                 275                 280                  285
Thr Thr Ser Ala Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Pro
                 290                 295                  300
Pro Ile Ser Ser Thr Ile Ser Val Gln Pro Ser Ser Ser Ser Ser
```

```
                305                 310                 315
Ser Pro Thr Thr Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Gly
                320                 325                 330
Ser Ala Pro Thr Thr Ser Ala Thr Ser Val Gln Pro Ser Ser Ser
                335                 340                 345
Ser Ser Pro Pro Ile Ser Ser Thr Ile Ser Val Gln Pro Ser Ser
                350                 355                 360
Ser Ser Ser Ser Pro Thr Thr Ser Thr Thr Ser Val Gln Pro Ser
                365                 370                 375
Ser Ser Gly Ser Ala Pro Thr Thr Ser Ala Thr Ser Val Gln Pro
                380                 385                 390
Ser Ser Ser Ser Ser Val Pro Thr Thr Ser Ala Thr Ser Val Arg
                395                 400                 405
Ser Ser Ser Ser Ser Ser Thr Pro Ile Pro Thr Thr Thr Ser Val
                410                 415                 420
Gln Pro Ser Ser Ser Ser Ser Val Pro Thr Thr Ser Ala Thr Ser
                425                 430                 435
Val Gln Thr Ser Ser Ser Ser Ser Thr Pro Ile Pro Ser Thr Thr
                440                 445                 450
Ser Val Gln Pro Ser Ser Ser Ser Ser Ala Pro Thr Thr Ser Ala
                455                 460                 465
Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Pro Pro Ile Ser Ser
                470                 475                 480
Thr Ile Ser Val Gln Pro Ser Ser Ser Ser Ser Ser Pro Thr Thr
                485                 490                 495
Ser Thr Thr Ser Val Gln Pro Ser Ser Ser Gly Ser Ala Pro Thr
                500                 505                 510
Thr Ser Ala Thr Ser Val Gln Pro Ser Ser Ser Ser Ser Pro Pro
                515                 520                 525
Ile Ser Ser
```